



Patent Application  
Attorney Docket No. PC10244A USA

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

RECEIVED

IN RE APPLICATION OF: :  
R. S. OBACH :  
APPLICATION NO.: 09/528798 :  
FILING DATE: March 21 2000 : DEC - 5 2003  
TITLE: USE OF A CYP2D6 INHIBITORS IN :  
COMBINATION THERAPIES :  
: :  
: :

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Commissioner for Patents  
P.O. Box 1450  
Alexandria, VA 22313-1450

**DECLARATION UNDER 37 C.F.R. 1.132**

Sir:

RONALD SCOTT OBACH, hereby declares, states and says that:

1. He received a B.S. from the State University of New York at Binghamton in 1985, and a Ph.D. from Brandeis University in 1990.
2. He is currently employed by Pfizer Inc. as a Research Advisor in the Pfizer research facility in Groton, Connecticut, and he has worked at Pfizer Inc. for 11 years.
3. He is familiar with the subject matter of the above-identified application and the references cited therein.
4. The above-identified application is directed to a method of administering the drug (2S,3S)-2-phenyl-3-(2-methoxy-5-trifluoromethoxyphenyl)methylamino-piperidine, or a pharmaceutically acceptable salt thereof, in combination with a CYP2D6 inhibitor, or a pharmaceutically acceptable salt thereof, to a human in need of the intended pharmaceutical activity of the drug, wherein the drug and the CYP2D6 inhibitor are not the same compound.

The CYP2D6 inhibitor may be, for example, quinidine, ajmalicine or pharmaceutically acceptable salts thereof.

5. In the enclosed data for the compound (2S,3S)-2-phenyl-3-(2-methoxy-5-trifluoromethoxyphenyl)methylamino-piperidine, denoted as "CP-B" in the data, Tables 1-4 describe enzymatic kinetic parameters for the metabolism of the compound (including O-demethylation and N-dealkylation) in various mammals, and Table 5 describes the inhibition of the same compound by Cytochrome P450 isoform specific inhibitors. In the figures, Figures 10 and 11 show a correlation between metabolism and inhibition of the same compound using quinidine (Figure 10) and ketoconazole (Figure 11).

6. The foregoing data and figures show a surprising effectiveness of (2S,3S)-2-phenyl-3-(2-methoxy-5-trifluoromethoxyphenyl)methylamino-piperidine in combination with a CYP2D6 inhibitor such as, for example, quinidine or ketoconazole.

He further declares that all statements made herein of his own knowledge are true and all statements made on information and belief are believed to be true. All statements made herein are made with the knowledge that willful false statements and the like so made are punishable by fine or imprisonment or both, under section 1001 of Title 18 of the United States Code, and that willful false statements may jeopardize the validity of the above application or any patent that may issue from it.

Date: 03-Dec-2003

  
\_\_\_\_\_  
Ronald Scott Obach

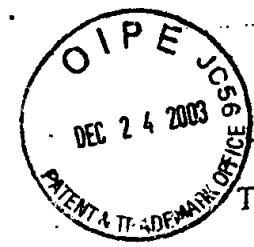


TABLE 1. ENZYME KINETIC PARAMETERS FOR METABOLISM OF  
CP-<sup>B</sup> IN POOLED HUMAN LIVER MICROSOMES

Parameter	(O-demethylation)	(N-dealkylation)
Kinetic Behavior	simple	sigmoidal
K <sub>Mapp</sub> ( $\mu$ M)	0.24	30
V <sub>max</sub> (pmol/min/mg)	14	150
CL' <sub>int</sub> ( $\mu$ L/min/mg)	59	5.2
Hill Coefficient	--	1.5
K <sub>M(free)</sub> ( $\mu$ M)	0.041	5.1
CL' <sub>int(free)</sub> ( $\mu$ L/min/mg)	350	31
scaled CL' <sub>int</sub> (mL/min/kg) <sup>a</sup>	53	4.7
scaled CL' <sub>int(free)</sub> (mL/min/kg) <sup>a</sup>	320	28

<sup>a</sup>Intrinsic clearance scaled per kg body weight using the values of 45 mg microsomal protein per gm liver and 20 gm liver per kg body weight in human.

"free" parameters are corrected for f<sub>u(microsomes)</sub> = 0.166

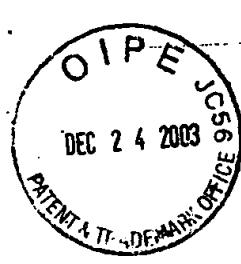
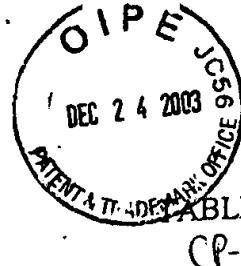


TABLE 2. ENZYME KINETIC PARAMETERS FOR METABOLISM OF  
CP. B IN POOLED RAT LIVER MICROSONES

Parameter	(O-demethylation)		(N-dealkylation)	
	Male	Female	Male	Female
Kinetic Behavior	sigmoidal/ substrate inhibition	sigmoidal/ substrate inhibition	biphasic	sigmoidal/ biphasic
$K_{Mapp}$ ( $\mu M$ )	0.44	0.36	43	88
$V_{max}$ (pmol/min/mg)	270	120	68	33
$CL'_{int(1)}$ ( $\mu L/min/mg$ )	610	330	1.6	0.40
$CL'_{int(2)}$ ( $\mu L/min/mg$ )	--	--	0.20	0.22
$CL'_{int(total)}$ ( $\mu L/min/mg$ )	610	330	1.8	0.62
$K_{iapp}$ ( $\mu M$ )	3.6	2.3	--	--
Hill Coefficient	1.5	1.7	--	4.3
$K_{M(free)}$ ( $\mu M$ )	0.070	0.047	6.9	11
$K_{iapp(free)}$ ( $\mu M$ )	0.58	0.30	--	--
$CL'_{int(total, free)}$ ( $\mu L/min/mg$ )	3800	2500	11	4.8
scaled $CL'_{int}$ ( $mL/min/kg$ ) <sup>a</sup>	1100	590	3.2	1.1
scaled $CL'_{int(free)}$ ( $mL/min/kg$ ) <sup>a</sup>	6800	4500	20	8.6

<sup>a</sup>Intrinsic clearance scaled per kg body weight using the values of 45 mg microsomal protein per gm liver and 40 gm liver per kg body weight in rat.

"free" parameters are corrected for  $f_{u(microsomes)} = 0.130$  for female, 0.160 for male



**TABLE 3. ENZYME KINETIC PARAMETERS FOR METABOLISM OF  
CP-B IN POOLED DOG LIVER MICROSOMES**

Parameter	(O-demethylation)	(N-dealkylation)	N-Hydroxylation
Kinetic Behavior	simple	sigmoidal	substrate inhibition
$K_{Mapp}$ ( $\mu M$ )	1.4	110	170
$V_{max}$ (pmol/min/mg)	210	240	880
$CL'_{int}$ ( $\mu L/min/mg$ )	140	2.2	5.2
$K_{iapp}$ ( $\mu M$ )	--	--	690
Hill Coefficient	--	1.5	--
$K_{M(free)}$ ( $\mu M$ )	0.20	15	24
$K_{iapp(free)}$ ( $\mu M$ )	--	--	97
$CL'_{int(free)}$ ( $\mu L/min/mg$ )	1000	16	37
scaled $CL'_{int}$ ( $mL/min/kg$ ) <sup>a</sup>	200	3.2	7.5
scaled $CL'_{int(free)}$ ( $mL/min/kg$ ) <sup>a</sup>	1400	23	53

<sup>a</sup>Intrinsic clearance scaled per kg body weight using the values of 45 mg microsomal protein per gm liver and 32 gm liver per kg body weight in dog.

"free" parameters are corrected for  $f_{u(microsomes)} = 0.144$

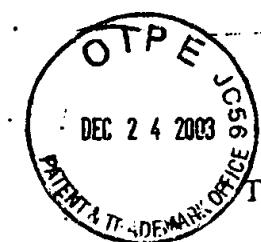


TABLE 4. ENZYME KINETIC PARAMETERS FOR METABOLISM OF  
CF-B<sub>1</sub> IN POOLED MONKEY LIVER MICROSOMES

Parameter	(O-demethylation)	(N-dealkylation)
Kinetic Behavior	simple	complex
K <sub>Mapp</sub> ( $\mu$ M)	0.73	80
V <sub>max</sub> (pmol/min/mg)	98	620
CL' <sub>int(1)</sub> ( $\mu$ L/min/mg)	130	7.7
CL' <sub>int(2)</sub> ( $\mu$ L/min/mg)	--	150
CL' <sub>int(total)</sub> ( $\mu$ L/min/mg)	130	160
K <sub>iapp</sub> ( $\mu$ M)	--	52
Hill Coefficient	--	1.1
K <sub>M(free)</sub> ( $\mu$ M)	0.15	16
K <sub>iapp(free)</sub> ( $\mu$ M)	--	10
CL' <sub>int(total, free)</sub> ( $\mu$ L/min/mg)	650	800
scaled CL' <sub>int</sub> (mL/min/kg) <sup>a</sup>	190	230
scaled CL' <sub>int(free)</sub> (mL/min/kg) <sup>a</sup>	940	1200

<sup>a</sup>Intrinsic clearance scaled per kg body weight using the values of 45 mg microsomal protein per gm liver and 32 gm liver per kg body weight in monkey.

"free" parameters are corrected for f<sub>u(microsomes)</sub> = 0.198



TABLE 5. INHIBITION OF HUMAN LIVER MICROSOMAL CP- $\beta$   
METABOLISM BY CYTOCHROME P450 ISOFORM SPECIFIC INHIBITORS

Inhibitor	(O-demethylation)	(N-dealkylation)
quinidine		
IC <sub>50</sub> ( $\mu$ M)	0.14	ND
maximum inhibition (%)	100	ND
ketoconazole		
IC <sub>50</sub> ( $\mu$ M)	ND	0.076
maximum inhibition (%)	ND	100
ND, not determined		

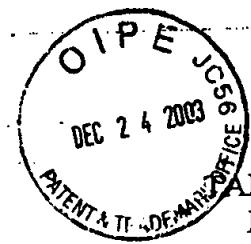


TABLE 6. CORRELATION OF CP<sub>B</sub> METABOLISM TO CYTOCHROME P450 SPECIFIC MARKER ACTIVITIES IN A PANEL OF HUMAN LIVER MICROSOMES

Cytochrome P450 Specific Marker Activity	Correlation Coefficient ( $r^2$ )	
	(O-demethylation)	(N-dealkylation)
phenacetin O-deethylase (CYP1A2)	0.003	0.002
tolbutamide hydroxylase (CYP2C9)	0.575	0.003
S-mephenytoin hydroxylase (CYP2C19)	0.094	0.074
bufuralol 1'-hydroxylase (CYP2D6)	0.863	0.001
testosterone 6 $\beta$ -hydroxylase (CYP3A)	0.071	0.870 <sup>a</sup>

a. Excludes one outlier point.

TABLE 7. METABOLISM OF CP-<sup>32</sup>B<sub>1</sub> BY HETEROLOGOUSLY EXPRESSED  
RECOMBINANT HUMAN CYTOCHROME P450 ENZYMES

CYP Enzyme	O-Demethylation	N-Dealkylation
	(pmol/min/nmol CYP) [S] = 0.2 μM	(pmol/min/nmol CYP) [S] = 25 μM
CYP1A1	37	480
CYP1A2	ND	ND
CYP2A6	ND	ND
CYP2B6	ND	ND
CYP2C9	ND	ND
CYP2C19	ND	ND
CYP2D6	4400	ND
CYP2E1	ND	ND
CYP3A4	ND	4500
CYP3A5	ND	9400

ND = None Detected

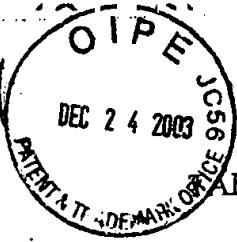


TABLE 8. ENZYME KINETIC PARAMETERS FOR CP- $\beta$ -METABOLISM  
BY HETEROLOGOUSLY EXPRESSED RECOMBINANT HUMAN  
CYTOCHROME P450 ENZYMES

Kinetic Behavior	rCYP2D6	rCYP3A4	rCYP3A5
	O-Demethylation	N-Dealkylation	N-Dealkylation
$K_{Mapp}$ ( $\mu M$ )	simple 0.057	substrate inhibition 11	substrate inhibition 22
$V_{max}$ (nmol/min/nmol CYP)	0.36	19	67
$CL'_{int}$ (mL/min/nmol CYP)	6.5	1.7	3.0
$K_{I(ap)}$ ( $\mu M$ )	--	1200	3800
$K_{M(free)}$ ( $\mu M$ )	0.041	1.4	3.7
$CL'_{int(free)}$ (mL/min/nmol CYP)	9.0	13	18
$K_{I(free)}$ ( $\mu M$ )	--	160	650

"free" parameters are corrected for  $f_{u(microsomes)} = 0.715, 0.133, \text{ and } 0.173$  for CYP2D6, 3A4, and 3A5, respectively.

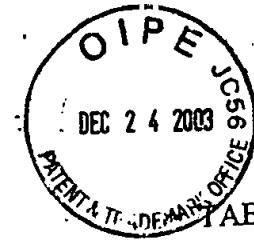


TABLE 9. NON-SPECIFIC BINDING OF CP-<sup>3</sup> TO LIVER MICROSONES AND RECOMBINANT CYP MICROSONES.

Replicate	Matrix	Non-specific Binding			Replicate	Matrix	Non-specific Binding	
		% Bound <sup>1</sup>	% Free				% Bound	% Free
1	HL-Mix-12	84.5	15.5	1	2	RL-137 (male)	84.9	15.1
	HL-Mix-12	82.3	17.7	2		RL-137 (male)	83.2	16.8
	Mean	83.4	16.6			Mean	84.1	16.0
	S.D.	1.6	1.6			S.D.	1.2	1.2
1	Dog Mix	83.2	16.8	1	2	rCYP3A4	86.8	13.2
	Dog Mix	88.0	12.0	2		rCYP3A4	86.7	13.3
	Mean	85.6	14.4			Mean	86.8	13.3
	S.D.	3.4	3.4			S.D.	0.1	0.1
1	Monkey Mix	80.7	19.3	1	2	rCYP3A5	83.4	16.6
	Monkey Mix	79.7	20.3	2		rCYP3A5	82.0	18.0
	Mean	80.2	19.8			Mean	82.7	17.3
	S.D.	0.7	0.7			S.D.	1.0	1.0
1	RL-129 (female)	88.2	11.8	1	2	rCYP2D6	25.7	74.3
	RL-129 (female)	85.9	14.1	2		rCYP2D6	31.4	68.6
	Mean	87.1	13.0			Mean	28.6	71.5
	S.D.	1.6	1.6			S.D.	4.0	4.0

<sup>1</sup> Non-specific binding calculated as the ratio of (microsomes - buffer) concentration divided by the microsome concentration

Microsomal Protein Concentration is 0.5 mg/mL for all human and animal species.  
 rCYP protein concentration is 2.0 mg/mL for CYP3A4 and 3A5, and 0.026 mg/mL for CYP2D6

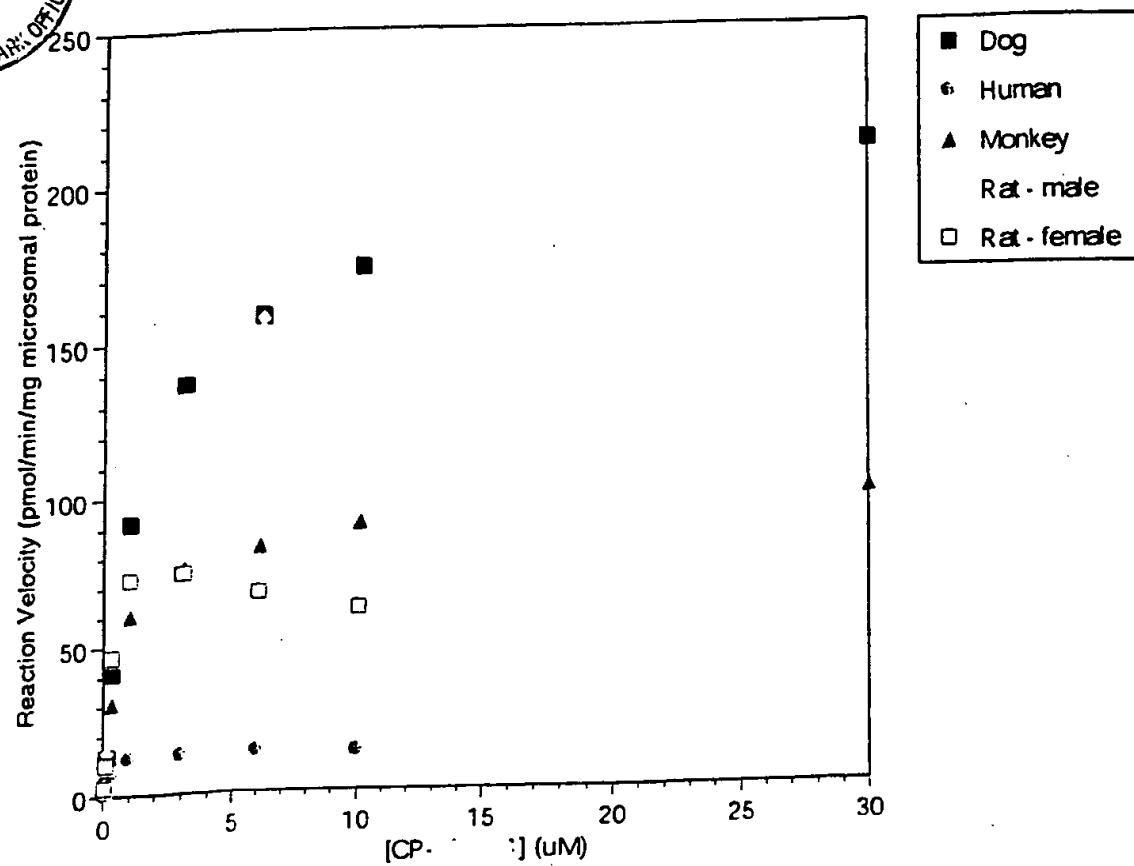
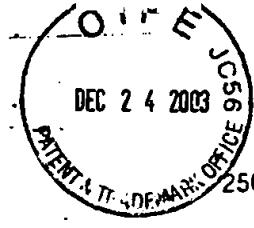


FIGURE 4. SUBSTRATE SATURATION CURVES FOR CP- $\beta$ -O-DEMETHYLATION IN LIVER MICROSOMES

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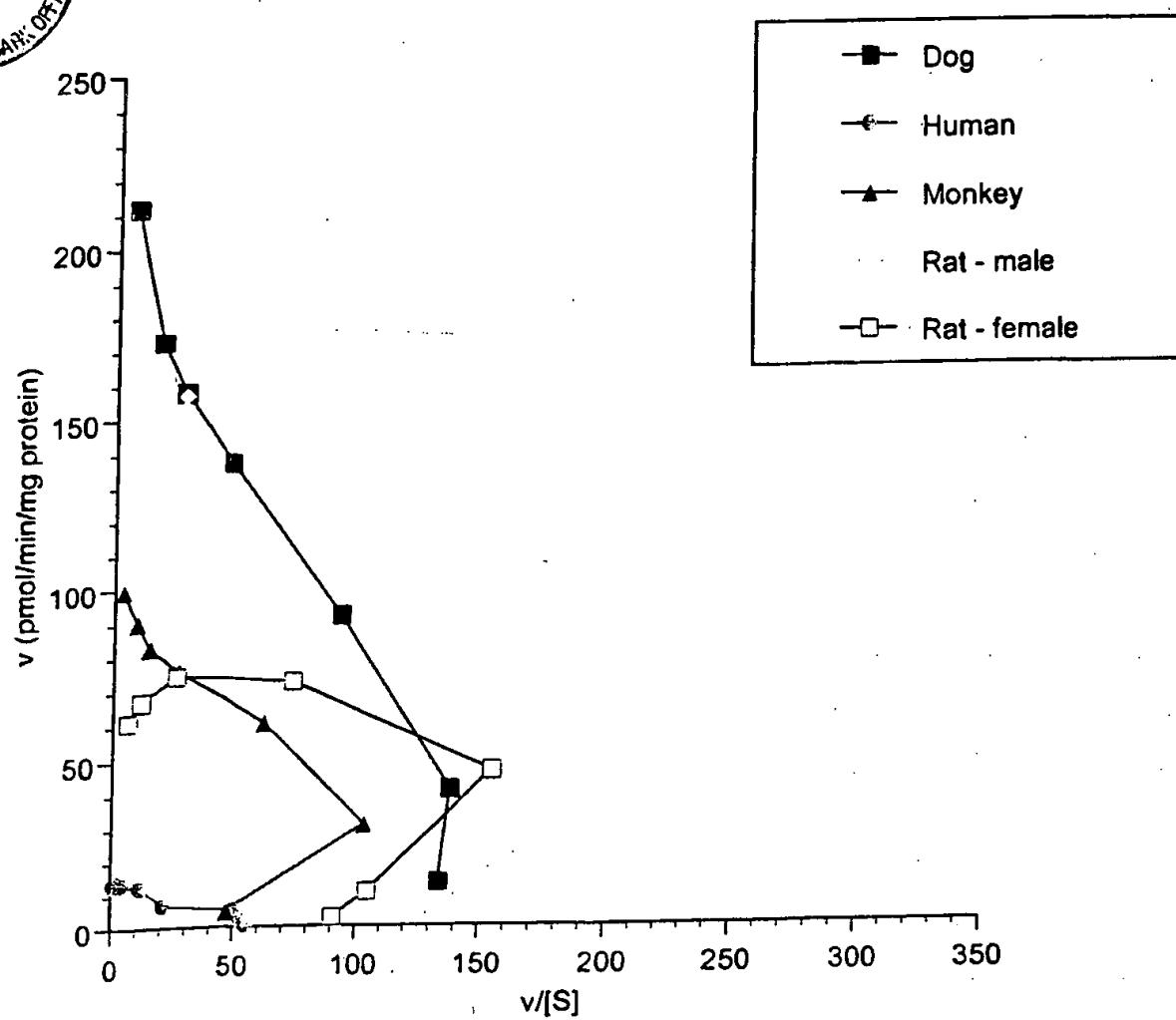


FIGURE 5. EADIE-HOFSTEE PLOT FOR CP-B O-DEMETHYLATION  
IN LIVER MICROSONES

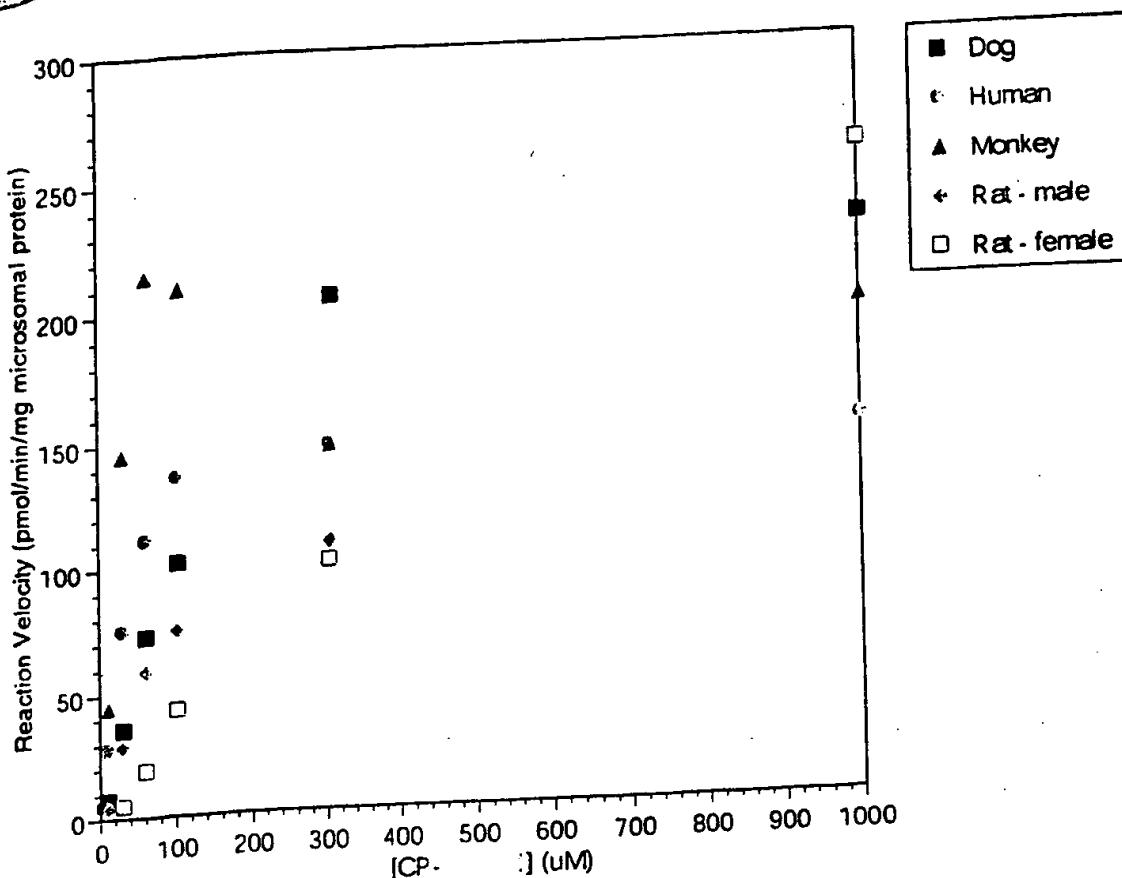


FIGURE 6. SUBSTRATE SATURATION CURVES FOR CP- $\beta$ -N-DEALKYLATION IN LIVER MICROSONES

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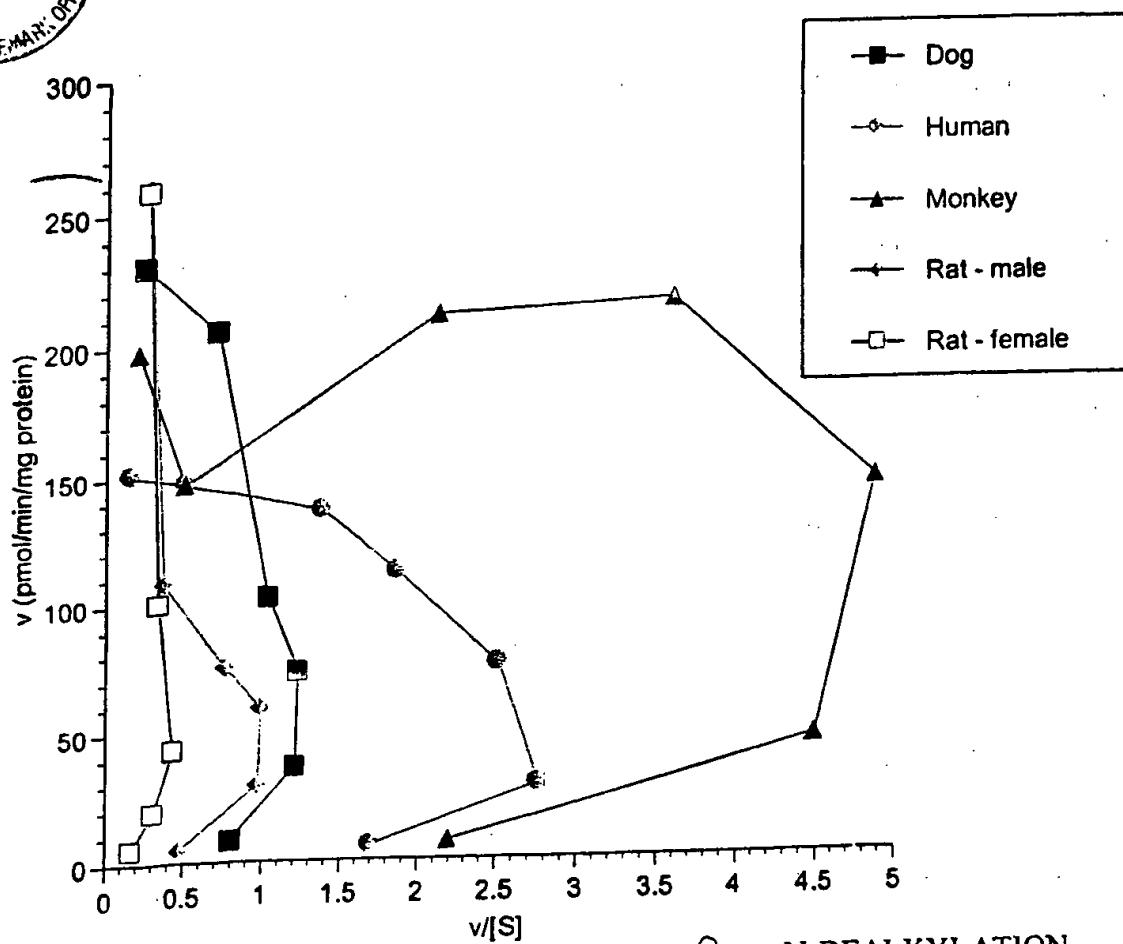


FIGURE 7. EADIE-HOFSTEE PLOT FOR CP- $\beta$ -N-DEALKYLATION KINETICS IN LIVER MICROSONES

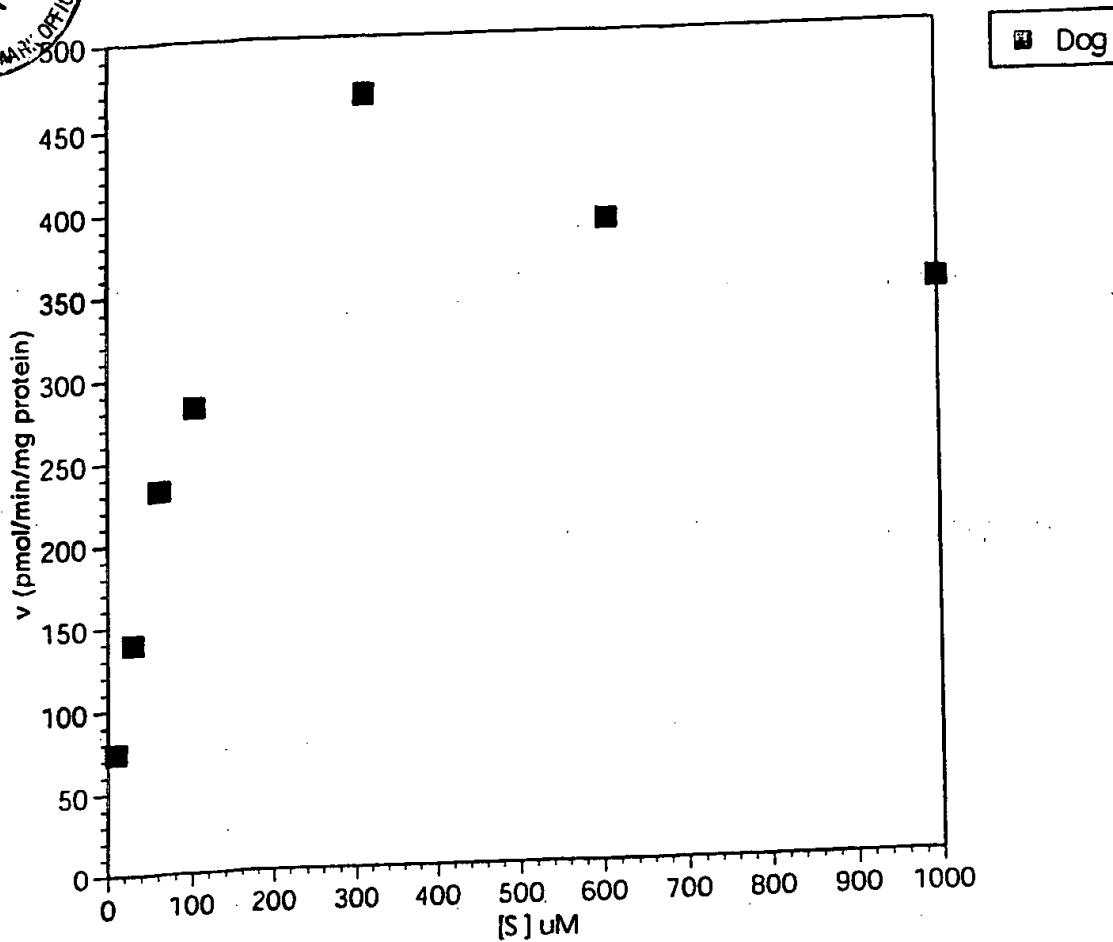
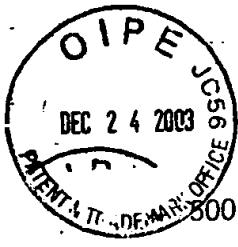


FIGURE 8. SUBSTRATE SATURATION CURVES FOR CP $\beta$  N-HYDROXYLATION IN DOG LIVER MICROSOMES

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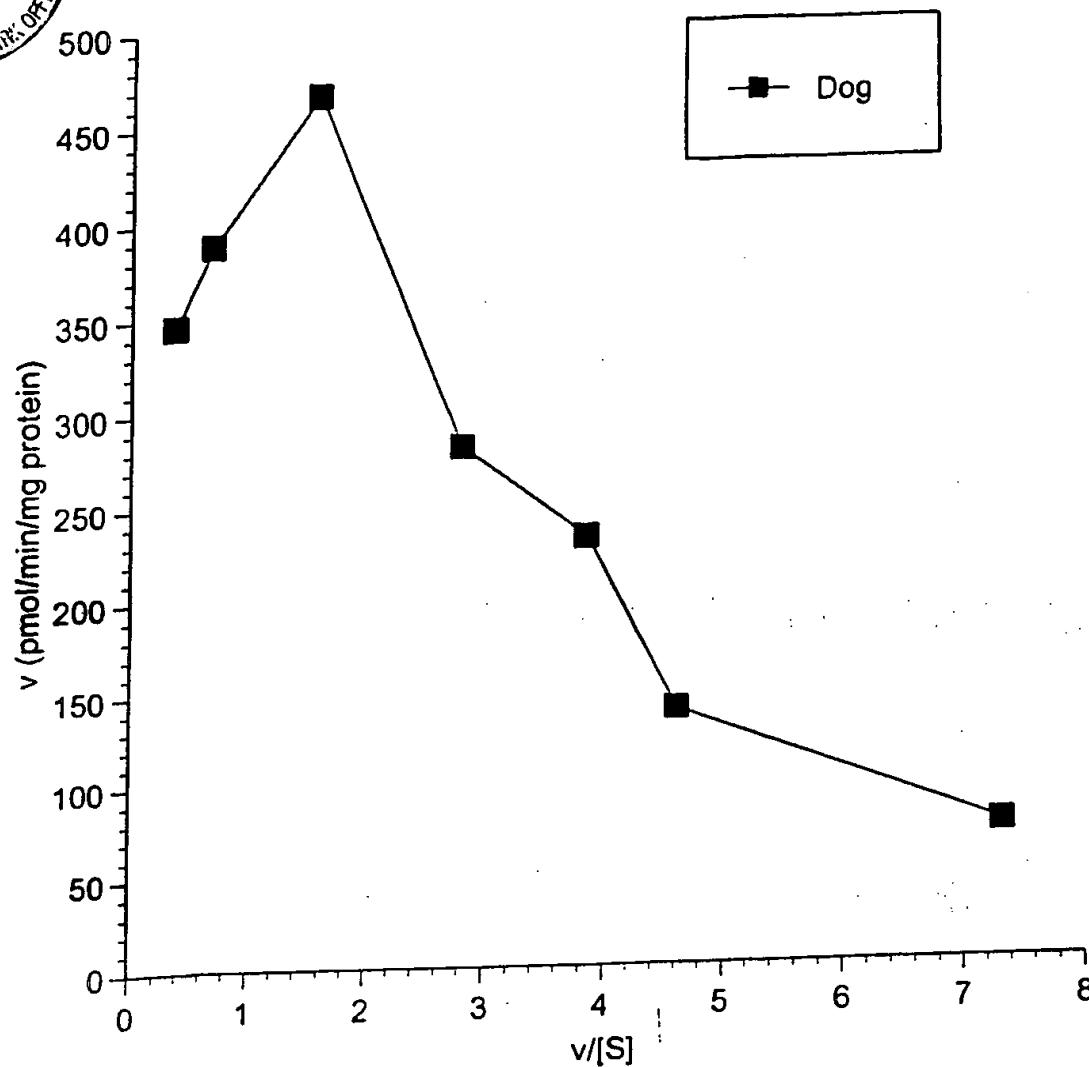


FIGURE 9. EADIE-HOFSTEE PLOT FOR CP- $\beta$  N-HYDROXYLATION  
KINETICS IN DOG LIVER MICROSONES

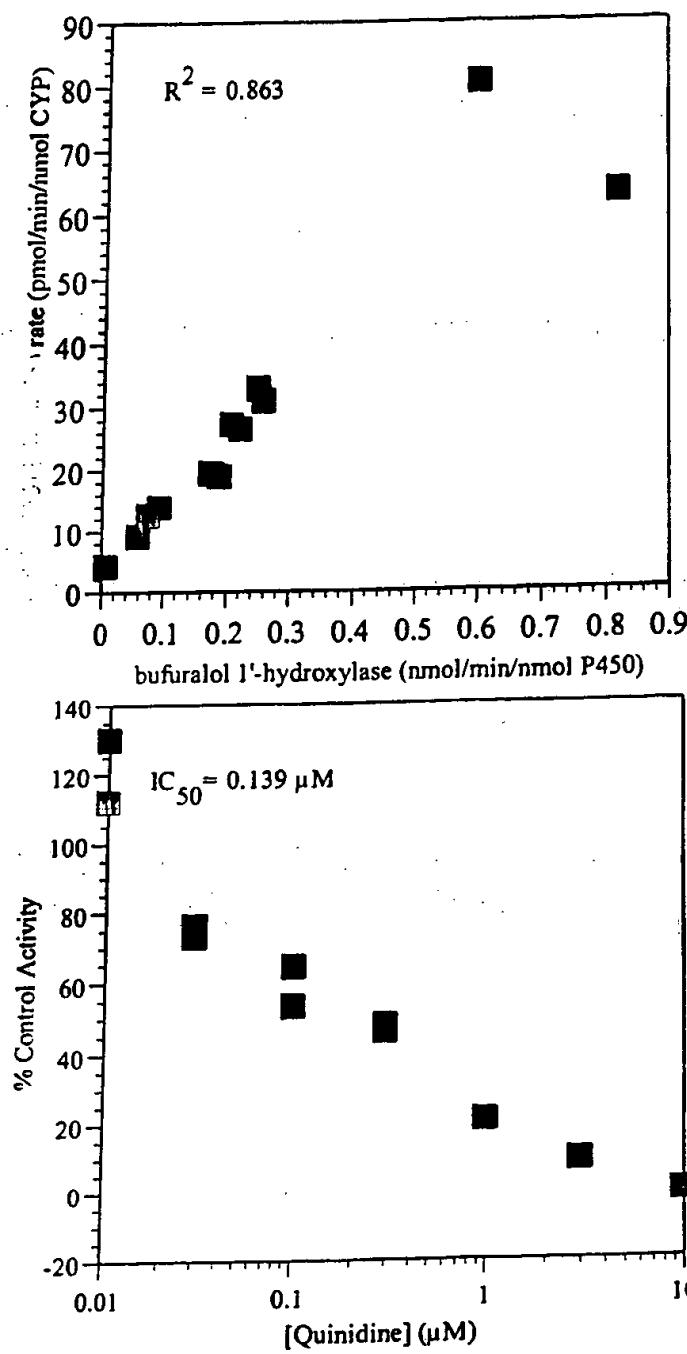
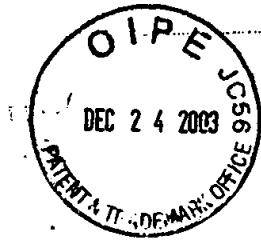


FIGURE 10. CP- $\beta$ -O-DEMETHYLATION: CORRELATION BETWEEN O-DEMETHYLATION AND BUFURALOL 1'-HYDROXYLASE ACTIVITIES IN HUMAN LIVER MICROSOMES (TOP) AND INHIBITION OF CP- $\beta$ -O-DEMETHYLATION USING QUINIDINE, A CYP2D6 SPECIFIC INHIBITOR IN HUMAN LIVER MICROSOMES (BOTTOM).

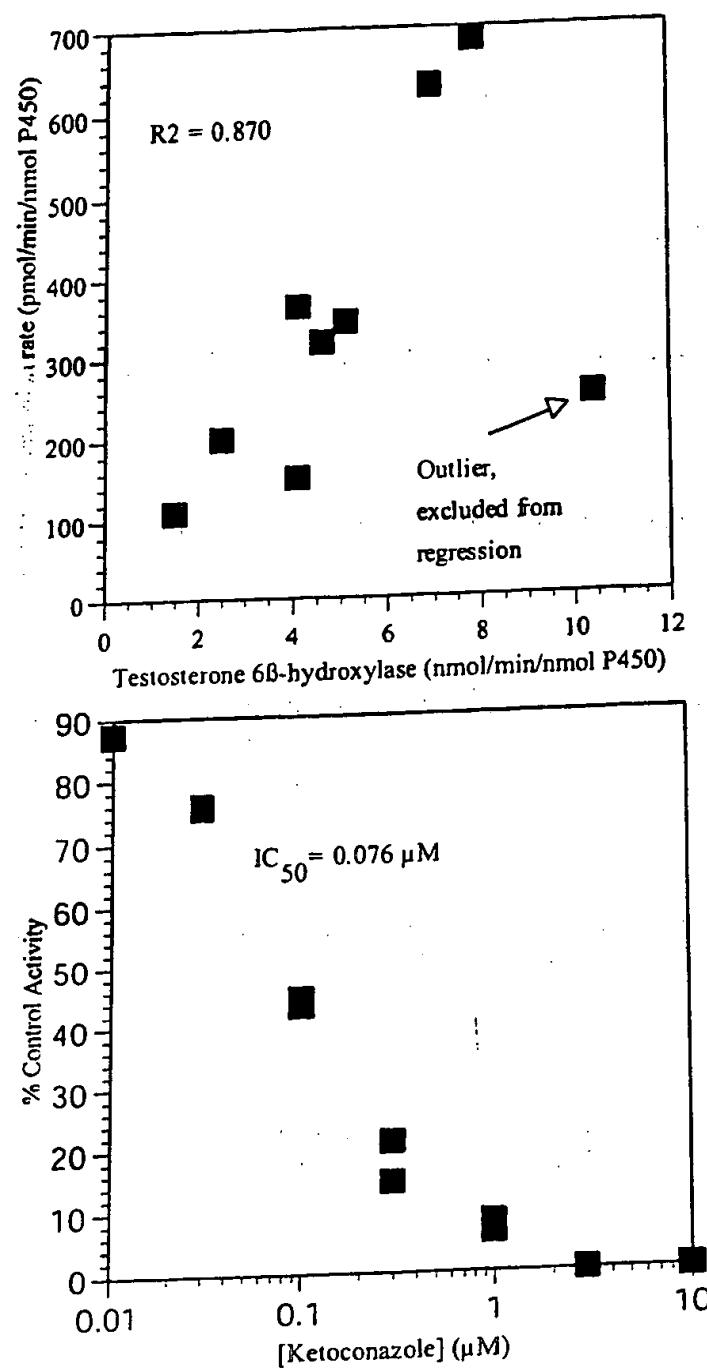
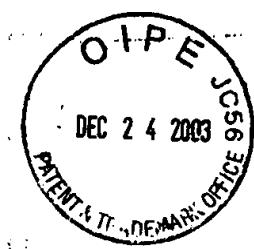


FIGURE 11. CP- $\beta$ -CP N-DEALKYLATION: CORRELATION BETWEEN N-DEALKYLATION AND TESTOSTERONE 6 $\beta$ -HYDROXYLASE ACTIVITIES IN HUMAN LIVER MICROSONES (TOP) AND INHIBITION OF CP- $\beta$ -CP N-DEALKYLATION USING KETOCONAZOLE (A CYP3A SELECTIVE INHIBITOR) IN HUMAN LIVER MICROSONES (BOTTOM).

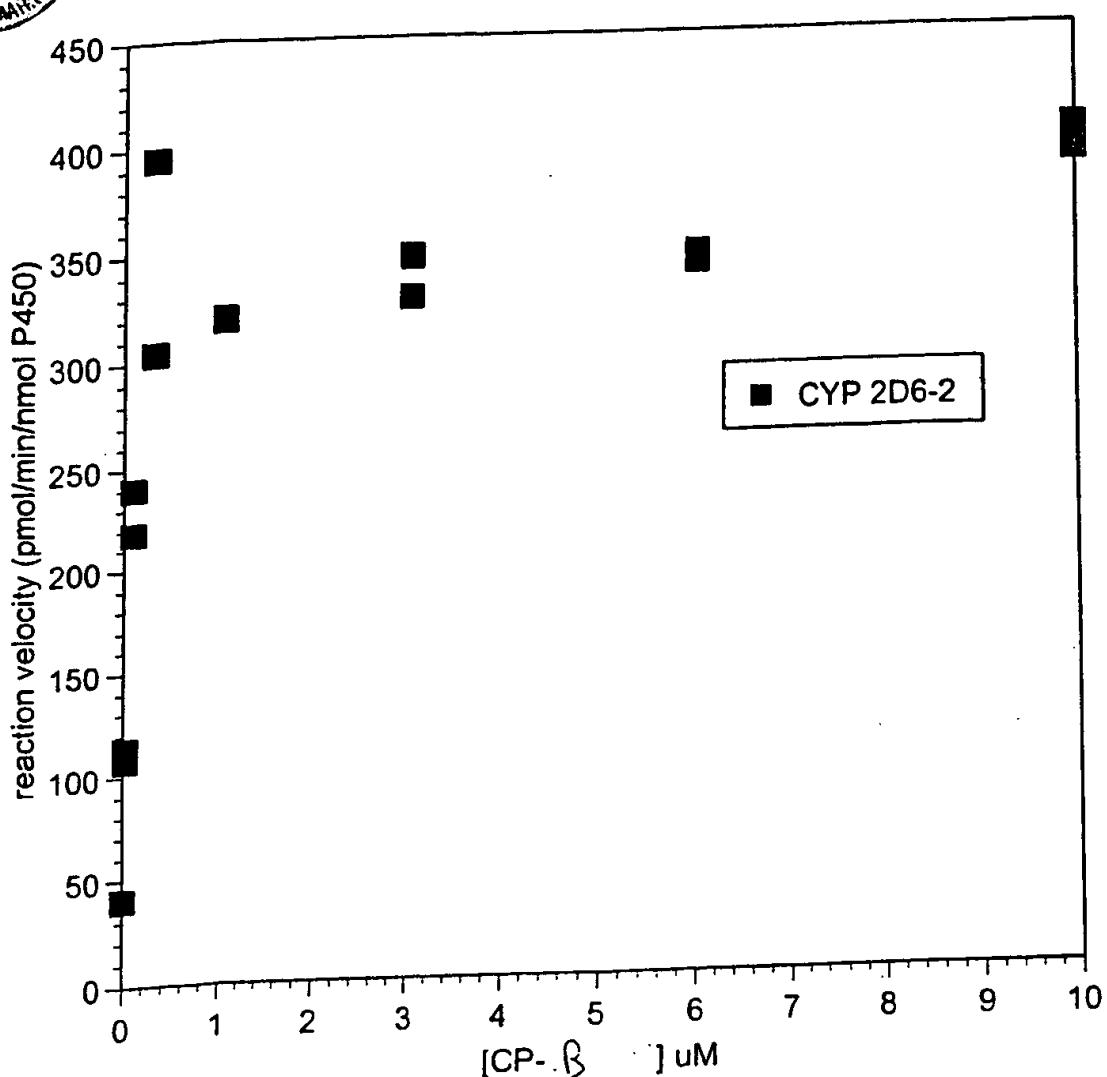
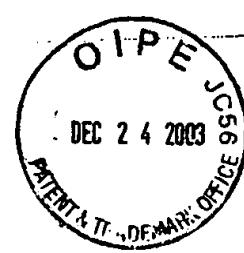


FIGURE 12. SUBSTRATE SATURATION PLOT OF CP- $\beta$   
O-DEMETHYLATION BY CYTOCHROME P4502D6

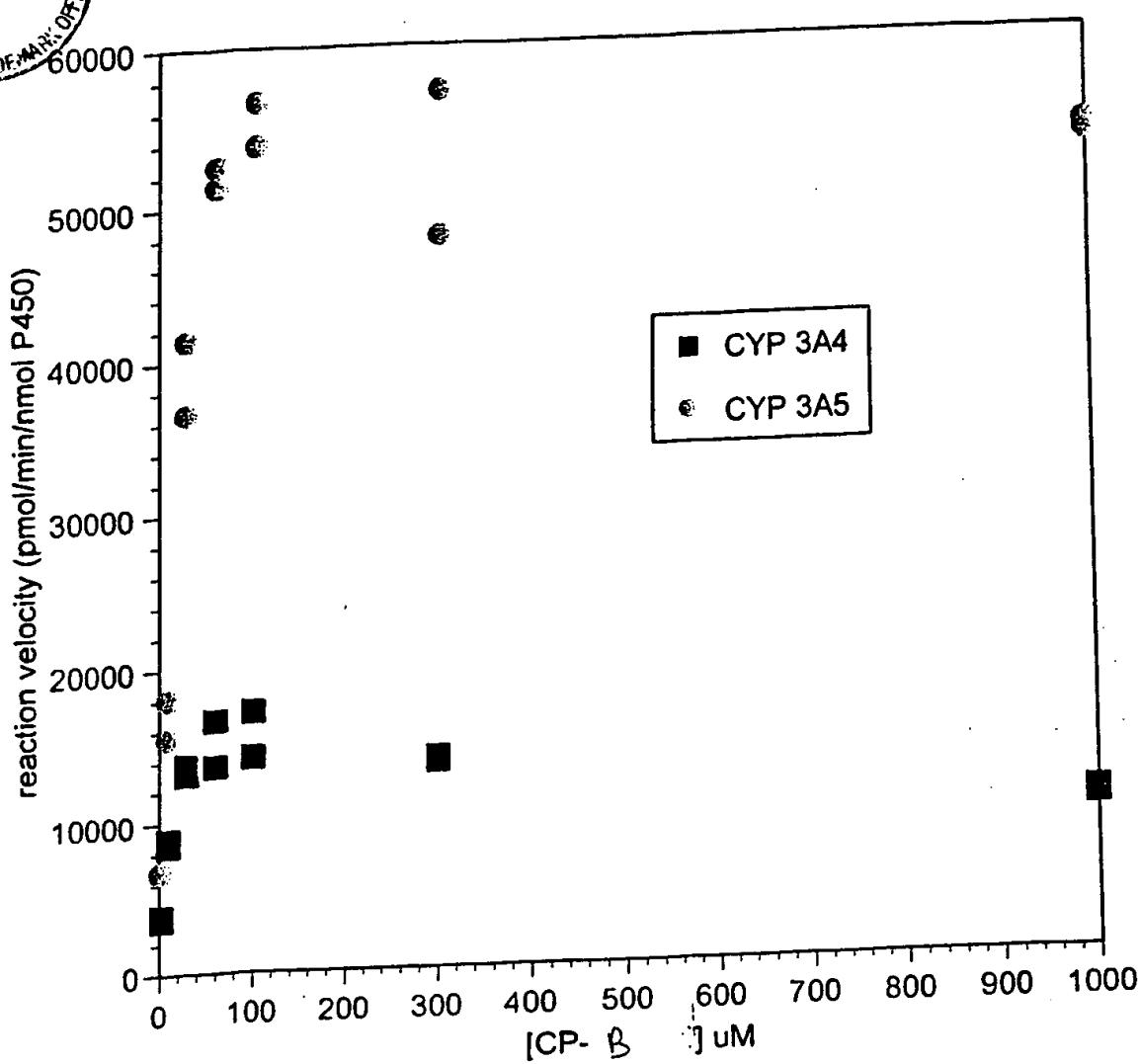


FIGURE 13. SUBSTRATE SATURATION PLOT OF CP- $\beta$   
N-DEALKYLATION BY CYTOCHROME P4503A4 AND 3A5